

## SEQUENCE LISTING

<110> Conklin, Darrell C.  
Haldeman, Betty A.  
Grossmann, Angelika

<120> MAMMALIAN CYTOKINE-LIKE POLYPEPTIDE-10

<130> 97-72

<160> 43

<170> FastSEQ for Windows Version 3.0

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<212> DNA

<213> Homo sapiens

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<221> CDS

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Ser Leu Ala Phe Ser Leu Leu Ser Ala Ala Phe Tyr Leu Leu Trp Thr	
5 10 15 20	

cct tcc act gga ctg aag aca ctc aat ttg gga agc tgt gtg atc gcc	152
Pro Ser Thr Gly Leu Lys Thr Leu Asn Leu Gly Ser Cys Val Ile Ala	
25 30 35	

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Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg Gly Ser	
40 45 50	

gtg caa gcc aaa gat gga aac att gac atc aga atc tta agg agg act	248
Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr	
55 60 65	

gag tct ttg caa gac aca aag cct gcg aat cga tgc tgc ctc ctg cgc 296  
 Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg  
       70                      75                      80

cat ttg cta aga ctc tat ctg gac agg gta ttt aaa aac tac cag acc 344  
 His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr  
       85                      90                      95                      100

cct gac cat tat act ctc cgg aag atc agc agc ctc gcc aat tcc ttt 392  
 Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe  
                       105                      110                      115

ctt acc atc aag aag gac ctc cgg ctc tgt cat gcc cac atg aca tgc 440  
 Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His Ala His Met Thr Cys  
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cat tgt ggg gag gaa gca atg aag aaa tac agc cag att ctg agt cac 488  
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                       135                      140                      145

ttt gaa aag ctg gaa cct cag gca gca gtt gtg aag gct ttg ggg gaa 536  
 Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu  
       150                      155                      160

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       165                      170                      175

gtgatgctgc tgctaagaat attcgaggtc aagagctcca gtcttcaata cctgcagagg 642  
 aggcatgacc ccaaaccacc atctctttac tgtactagtc ttgtgctggc cacagtgtat 702  
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 gagaccatac ttgtataaga tttttgtaat atctttctgc tattggatat atttattagt 822  
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			20					25					30		
Cys	Val	Ile	Ala	Thr	Asn	Leu	Gln	Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp
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Ile	Arg	Gly	Ser	Val	Gln	Ala	Lys	Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile
	50					55					60				
Leu	Arg	Arg	Thr	Glu	Ser	Leu	Gln	Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys
65					70					75					80
Cys	Leu	Leu	Arg	His	Leu	Leu	Arg	Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys
				85					90					95	
Asn	Tyr	Gln	Thr	Pro	Asp	His	Tyr	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
			100					105					110		
Ala	Asn	Ser	Phe	Leu	Thr	Ile	Lys	Lys	Asp	Leu	Arg	Leu	Cys	His	Ala
		115					120					125			
His	Met	Thr	Cys	His	Cys	Gly	Glu	Glu	Ala	Met	Lys	Lys	Tyr	Ser	Gln
	130					135					140				
Ile	Leu	Ser	His	Phe	Glu	Lys	Leu	Glu	Pro	Gln	Ala	Ala	Val	Val	Lys
145					150					155					160
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 5 10 15 20

cct tcc act gga ctg aag aca ctc aat ttg gga agc tgt gtg atc gcc 152  
Pro Ser Thr Gly Leu Lys Thr Leu Asn Leu Gly Ser Cys Val Ile Ala  
25 30 35

aca aac ctt cag gaa ata cga aat gga ttt tct gac ata cgg ggc agt	200
Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg Gly Ser	
40 45 50	
gtg caa gcc aaa gat gga aac att gac atc aga atc tta agg agg act	248
Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr	
55 60 65	
gag tct ttg caa gac aca aag cct gcg aat cga tgc tgc ctc ctg cgc	296
Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg	
70 75 80	
cat ttg cta aga ctc tat ctg gac agg gta ttt aaa aac tac cag acc	344
His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr	
85 90 95 100	
cct gac cat tat act ctc cgg aag atc agc agc ctc gcc aat tcc ttt	392
Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe	
105 110 115	
ctt acc atc aag aag gac ctc cgg ctc tgt ctg gaa cct cag gca gca	440
Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys Leu Glu Pro Gln Ala Ala	
120 125 130	
gtt gtg aag gct ttg ggg gaa cta gac att ctt ctg caa tgg atg gag	488
Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met Glu	
135 140 145	
gag aca gaa taggaggaaa gtgatgctgc tgctaagaat attcgaggtc	537
Glu Thr Glu	
150	
aagagctcca gtcttcaata cctgcagagg aggcatgacc ccaaaccacc atctctttac	597
tgtactagtc ttgtgctggt cacagtgtat cttattttatg cattacttgc ttccttgc	657
gattgtcttt atgcatcccc aatcttaatt gagaccatac ttgtataaga tttttgta	717
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&lt;210&gt; 4

&lt;211&gt; 151

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 4

Met Lys Ala Ser Ser Leu Ala Phe Ser Leu Leu Ser Ala Ala Phe Tyr  
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 Leu Leu Trp Thr Pro Ser Thr Gly Leu Lys Thr Leu Asn Leu Gly Ser  
 20 25 30  
 Cys Val Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp  
 35 40 45  
 Ile Arg Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile  
 50 55 60  
 Leu Arg Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys  
 65 70 75 80  
 Cys Leu Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys  
 85 90 95  
 Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu  
 100 105 110  
 Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys Leu Glu  
 115 120 125  
 Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu  
 130 135 140  
 Gln Trp Met Glu Glu Thr Glu  
 145 150

&lt;210&gt; 5

&lt;211&gt; 253

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 5

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 tttctgagat acggggcagt gtgcaagcca aagatggaaa cattgacatc agaatcttaa 240  
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&lt;210&gt; 6

&lt;211&gt; 24

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 6

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24

&lt;210&gt; 7

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 ttgctaagac tctatctgga cagggtatth aaaaactacc agaccctga ccattatact 180  
 ctccggaaga tcagcagcct cgccaattcc tttcttacca tcaagaagga cctccggctc 240  
 tgtcatgccc acatgacatg ccattgtggg gaggaagcaa tgaagaaata cagccagatt 300  
 ctgagtcact ttgaaaagct ggaacctcag gcagcagttg tgaaggcttt gggggaacta 360  
 gacattcttc tgcaatggat ggaggagaca gaataggagg aaagtgatgc tgctgctaag 420  
 aatattcgag gtcaagagct ccagtcttca atacctgcag aggaggcatg accccaaacc 480  
 accatctctt tactgtacta gtcttgtgct ggtcacagtg tatcttattt atgcattact 540  
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 agatthttgt aatatctttc tgctattgga tatatttatt agttaatata tttatttatt 660  
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 ttgctaagac tctatctgga cagggtattt aaaaactacc agaccctga ccattatact 180  
 ctccggaaga tcagcagcct cgccaattcc tttcttacca tcaagaagga cctccggctc 240  
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 Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg Arg Thr Glu Ser Leu Gln  
 35 40 45  
 Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu Leu Arg His Leu Leu Arg  
 50 55 60  
 Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr  
 65 70 75 80  
 Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys  
 85 90 95  
 Lys Asp Leu Arg Leu Cys His Ala His Met Thr Cys His Cys Gly Glu  
 100 105 110  
 Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe Glu Lys Leu  
 115 120 125  
 Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu  
 130 135 140  
 Leu Gln Trp Met Glu Glu Thr Glu

145

150

&lt;210&gt; 13

&lt;211&gt; 127

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 13

Leu	Lys	Thr	Leu	Asn	Leu	Gly	Ser	Cys	Val	Ile	Ala	Thr	Asn	Leu	Gln
1				5					10					15	



Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp	Ile	Arg	Gly	Ser	Val	Gln	Ala	Lys
			20					25					30		
Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile	Leu	Arg	Arg	Thr	Glu	Ser	Leu	Gln
		35					40					45			
Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys	Cys	Leu	Leu	Arg	His	Leu	Leu	Arg
		50				55					60				
Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys	Asn	Tyr	Gln	Thr	Pro	Asp	His	Tyr
65					70					75					80
Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu	Ala	Asn	Ser	Phe	Leu	Thr	Ile	Lys
				85					90					95	
Lys	Asp	Leu	Arg	Leu	Cys	Leu	Glu	Pro	Gln	Ala	Ala	Val	Val	Lys	Ala
			100					105					110		
Leu	Gly	Glu	Leu	Asp	Ile	Leu	Leu	Gln	Trp	Met	Glu	Glu	Thr	Glu	
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 <213> Homo sapiens

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 <213> Homo sapiens

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 <213> Homo sapiens

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<210> 18  
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 Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala  
 1 5 10  
 gtg ggt ttt ctt ctc tgg act cct tta act ggg ctc aag acc ctc cat 157  
 Val Gly Phe Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His  
 15 20 25  
 ttg gga agc tgt gtg att act gca aac cta cag gca ata caa aag gaa 205  
 Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu  
 30 35 40 45  
 ttt tct gag att cgg gat agt gtg caa gct gaa gat aca aat att gac 253  
 Phe Ser Glu Ile Arg Asp Ser Val Gln Ala Glu Asp Thr Asn Ile Asp  
 50 55 60  
 atc aga att tta agg acg act gag tct ttg aaa gac ata aag tct ttg 301  
 Ile Arg Ile Leu Arg Thr Thr Glu Ser Leu Lys Asp Ile Lys Ser Leu  
 65 70 75  
 gat agg tgc tgc ttc ctt cgt cat cta gtg aga ttc tat ctg gac agg 349  
 Asp Arg Cys Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg  
 80 85 90

gta ttc aaa gtc tac cag acc cct gac cac cat acc ctg aga aag atc 397  
Val Phe Lys Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile  
95 100 105

agc agc ctc gcc aac tcc ttt ctt atc atc aag aag gac ctc tca gtc 445  
Ser Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val  
110 115 120 125

tgt cat tct cac atg gca tgt cat tgt ggg gaa gaa gca atg gag aaa 493  
Cys His Ser His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys  
130 135 140

tac aac caa att ctg agt cac ttc ata gag ttg gaa ctt cag gca gcg 541  
Tyr Asn Gln Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala  
145 150 155

gtg gta aag gct ttg gga gaa cta ggc att ctt ctg aga tgg atg gag 589  
Val Val Lys Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu  
160 165 170

gag atg cta tagatgaaag tggagaggct gctgagaaca ctctgtcca 638  
Glu Met Leu  
175

agaatctcag	acctcagcac	catgaagaca	tggccccagg	tgtctggcatt	tctactcaag	698
agttccagtc	ctcagcacca	cgaagatggc	ctcaaaccac	caccctttg	tgatataact	758
tagtgctagc	tatgtgtata	ttattttctac	attattggct	cccttatgtg	aatgccttca	818
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<210> 19

<211> 176

&lt;212&gt; PRT

<213> Mus musculus

<400> 19

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			20					25					30		
Cys	Val	Ile	Thr	Ala	Asn	Leu	Gln	Ala	Ile	Gln	Lys	Glu	Phe	Ser	Glu
		35					40					45			
Ile	Arg	Asp	Ser	Val	Gln	Ala	Glu	Asp	Thr	Asn	Ile	Asp	Ile	Arg	Ile
	50					55					60				

Leu Arg Thr Thr Glu Ser Leu Lys Asp Ile Lys Ser Leu Asp Arg Cys  
 65 70 75 80  
 Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys  
 85 90 95  
 Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu  
 100 105 110  
 Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser  
 115 120 125  
 His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln  
 130 135 140  
 Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys  
 145 150 155 160  
 Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu  
 165 170 175

<210> 20  
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 Asp Thr Asn Ile Asp Ile Arg Ile Leu Arg Thr Thr Glu Ser Leu Lys  
 35 40 45  
 Asp Ile Lys Ser Leu Asp Arg Cys Cys Phe Leu Arg His Leu Val Arg  
 50 55 60  
 Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln Thr Pro Asp His His  
 65 70 75 80  
 Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys  
 85 90 95  
 Lys Asp Leu Ser Val Cys His Ser His Met Ala Cys His Cys Gly Glu  
 100 105 110  
 Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser His Phe Ile Glu Leu  
 115 120 125  
 Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Gly Ile Leu  
 130 135 140  
 Leu Arg Trp Met Glu Glu Met Leu  
 145 150

<210> 21

<211> 16  
 <212> PRT  
 <213> Mus musculus

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 1 5 10 15

<210> 25  
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 <212> PRT  
 <213> Mus musculus

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 1 5 10 15

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			20					25					30		
Leu	Arg	Thr	Thr	Glu	Ser	Leu	Lys	Asp	Ile	Lys	Ser	Leu	Asp	Arg	Cys
		35					40					45			
Cys	Phe	Leu	Arg	His	Leu	Val	Arg	Phe	Tyr	Leu	Asp	Arg	Val	Phe	Lys
	50					55					60				
Val	Tyr	Gln	Thr	Pro	Asp	His	His	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
65					70					75					80
Ala	Asn	Ser	Phe	Leu	Ile	Ile	Lys	Lys	Asp	Leu	Ser	Val	Cys	His	Ser
				85					90					95	
His	Met	Ala	Cys	His	Cys	Gly	Glu	Glu	Ala	Met	Glu	Lys	Tyr	Asn	Gln
			100					105					110		
Ile	Leu	Ser	His	Phe	Ile	Glu	Leu	Glu	Leu	Gln	Ala	Ala	Val	Val	Lys
		115					120					125			
Ala	Leu	Gly	Glu	Leu	Gly	Ile	Leu	Leu	Arg	Trp	Met	Glu	Glu	Met	Leu
	130					135						140			

&lt;210&gt; 26

&lt;211&gt; 144

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 26

Cys	Val	Ile	Ala	Thr	Asn	Leu	Gln	Glu	Ile	Arg	Asn	Gly	Phe	Ser	Asp
1				5					10					15	
Ile	Arg	Gly	Ser	Val	Gln	Ala	Lys	Asp	Gly	Asn	Ile	Asp	Ile	Arg	Ile
			20					25					30		
Leu	Arg	Arg	Thr	Glu	Ser	Leu	Gln	Asp	Thr	Lys	Pro	Ala	Asn	Arg	Cys
		35					40					45			
Cys	Leu	Leu	Arg	His	Leu	Leu	Arg	Leu	Tyr	Leu	Asp	Arg	Val	Phe	Lys
	50					55					60				
Asn	Tyr	Gln	Thr	Pro	Asp	His	Tyr	Thr	Leu	Arg	Lys	Ile	Ser	Ser	Leu
65					70					75					80
Ala	Asn	Ser	Phe	Leu	Thr	Ile	Lys	Lys	Asp	Leu	Arg	Leu	Cys	His	Ala
				85					90					95	
His	Met	Thr	Cys	His	Cys	Gly	Glu	Glu	Ala	Met	Lys	Lys	Tyr	Ser	Gln
			100					105					110		
Ile	Leu	Ser	His	Phe	Glu	Lys	Leu	Glu	Pro	Gln	Ala	Ala	Val	Val	Lys
		115					120					125			
Ala	Leu	Gly	Glu	Leu	Asp	Ile	Leu	Leu	Gln	Trp	Met	Glu	Glu	Thr	Glu
	130					135						140			

&lt;210&gt; 27

<211> 38  
 <212> PRT  
 <213> Homo sapiens

<400> 27  
 Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe  
 1 5 10 15  
 Glu Lys Leu Glu Pro Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu  
 20 25 30  
 Asp Ile Leu Leu Gln Trp  
 35

<210> 28  
 <211> 71  
 <212> PRT  
 <213> Homo sapiens

<400> 28  
 Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg  
 1 5 10 15  
 Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg  
 20 25 30  
 Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu  
 35 40 45  
 Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr  
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 Gln Thr Pro Asp His Tyr Thr  
 65 70

<210> 29  
 <211> 92  
 <212> PRT  
 <213> Homo sapiens

<400> 29  
 Ile Ala Thr Asn Leu Gln Glu Ile Arg Asn Gly Phe Ser Asp Ile Arg  
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 Gly Ser Val Gln Ala Lys Asp Gly Asn Ile Asp Ile Arg Ile Leu Arg  
 20 25 30  
 Arg Thr Glu Ser Leu Gln Asp Thr Lys Pro Ala Asn Arg Cys Cys Leu  
 35 40 45  
 Leu Arg His Leu Leu Arg Leu Tyr Leu Asp Arg Val Phe Lys Asn Tyr  
 50 55 60

Gln Thr Pro Asp His Tyr Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn  
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 Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys  
 85 90

<210> 30  
 <211> 82  
 <212> PRT  
 <213> Homo sapiens

<400> 30  
 Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu  
 1 5 10 15  
 Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp  
 20 25 30  
 Leu Arg Leu Cys His Ala His Met Thr Cys His Cys Gly Glu Glu Ala  
 35 40 45  
 Met Lys Lys Tyr Ser Gln Ile Leu Ser His Phe Glu Lys Leu Glu Pro  
 50 55 60  
 Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln  
 65 70 75 80  
 Trp Met

<210> 31  
 <211> 36  
 <212> PRT  
 <213> Homo sapiens

<400> 31  
 Leu Asp Arg Val Phe Lys Asn Tyr Gln Thr Pro Asp His Tyr Thr Leu  
 1 5 10 15  
 Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp  
 20 25 30  
 Leu Arg Leu Cys  
 35

<210> 32  
 <211> 61  
 <212> PRT  
 <213> Homo sapiens

<400> 32



Leu Ala Asn Ser Phe Leu Thr Ile Lys Lys Asp Leu Arg Leu Cys His  
 1 5 10 15  
 Ala His Met Thr Cys His Cys Gly Glu Glu Ala Met Lys Lys Tyr Ser  
 20 25 30  
 Gln Ile Leu Ser His Phe Glu Lys Leu Glu Pro Gln Ala Ala Val Val  
 35 40 45  
 Lys Ala Leu Gly Glu Leu Asp Ile Leu Leu Gln Trp Met  
 50 55 60

<210> 33  
 <211> 756  
 <212> DNA  
 <213> Mus musculus

<220>  
 <221> CDS  
 <222> (71)...(532)

<400> 33

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 Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala  
 1 5 10

gtg ggt ttt ctt ctc tgg act cct tta act ggg ctc aag acc ctc cat 157  
 Val Gly Phe Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His  
 15 20 25

ttg gga agc tgt gtg att act gca aac cta cag gca ata caa aag gaa 205  
 Leu Gly Ser Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu  
 30 35 40 45

ttt tct gag att cgg gat agt gtg tct ttg gat agg tgc tgc ttc ctt 253  
 Phe Ser Glu Ile Arg Asp Ser Val Ser Leu Asp Arg Cys Cys Phe Leu  
 50 55 60

cgt cat cta gtg aga ttc tat ctg gac agg gta ttc aaa gtc tac cag 301  
 Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln  
 65 70 75

acc cct gac cac cat acc ctg aga aag atc agc agc ctc gcc aac tcc 349  
 Thr Pro Asp His His Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser  
 80 85 90

ttt ctt atc atc aag aag gac ctc tca gtc tgt cat tct cac atg gca	397
Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys His Ser His Met Ala	
95 100 105	
tgt cat tgt ggg gaa gaa gca atg gag aaa tac aac caa att ctg agt	445
Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser	
110 115 120 125	
cac ttc ata gag ttg gaa ctt cag gca gcg gtg gta aag gct ttg gga	493
His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly	
130 135 140	
gaa cta ggc att ctt ctg aga tgg atg gag gag atg cta tagatgaaag	542
Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu Met Leu	
145 150	
tggataggct gctgagaaca ctctgtcca agaatctcag acctcagcac catgaagaca	602
tggccccagg tgctggcatt tctactcaag agttccagtc ctccagcacca cgaagatggc	662
ctcaaaccac caccctttg tgatataact tagtgctagc tatgtgtata ttatttctac	722
attattggct cccttatgtg aatgccttca tgtg	756

&lt;210&gt; 34

&lt;211&gt; 154

&lt;212&gt; PRT

&lt;213&gt; Mus musculus

&lt;400&gt; 34

Met Lys Gly Phe Gly Leu Ala Phe Gly Leu Phe Ser Ala Val Gly Phe	
1 5 10 15	
Leu Leu Trp Thr Pro Leu Thr Gly Leu Lys Thr Leu His Leu Gly Ser	
20 25 30	
Cys Val Ile Thr Ala Asn Leu Gln Ala Ile Gln Lys Glu Phe Ser Glu	
35 40 45	
Ile Arg Asp Ser Val Ser Leu Asp Arg Cys Cys Phe Leu Arg His Leu	
50 55 60	
Val Arg Phe Tyr Leu Asp Arg Val Phe Lys Val Tyr Gln Thr Pro Asp	
65 70 75 80	
His His Thr Leu Arg Lys Ile Ser Ser Leu Ala Asn Ser Phe Leu Ile	
85 90 95	
Ile Lys Lys Asp Leu Ser Val Cys His Ser His Met Ala Cys His Cys	
100 105 110	
Gly Glu Glu Ala Met Glu Lys Tyr Asn Gln Ile Leu Ser His Phe Ile	

115                      120                      125  
 Glu Leu Glu Leu Gln Ala Ala Val Val Lys Ala Leu Gly Glu Leu Gly  
 130                      135                      140  
 Ile Leu Leu Arg Trp Met Glu Glu Met Leu  
 145                      150

<210> 35  
 <211> 130  
 <212> PRT  
 <213> Mus musculus

<400> 35  
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 1                      5                      10                      15  
 Ala Ile Gln Lys Glu Phe Ser Glu Ile Arg Asp Ser Val Ser Leu Asp  
 20                      25                      30  
 Arg Cys Cys Phe Leu Arg His Leu Val Arg Phe Tyr Leu Asp Arg Val  
 35                      40                      45  
 Phe Lys Val Tyr Gln Thr Pro Asp His His Thr Leu Arg Lys Ile Ser  
 50                      55                      60  
 Ser Leu Ala Asn Ser Phe Leu Ile Ile Lys Lys Asp Leu Ser Val Cys  
 65                      70                      75                      80  
 His Ser His Met Ala Cys His Cys Gly Glu Glu Ala Met Glu Lys Tyr  
 85                      90                      95  
 Asn Gln Ile Leu Ser His Phe Ile Glu Leu Glu Leu Gln Ala Ala Val  
 100                      105                      110  
 Val Lys Ala Leu Gly Glu Leu Gly Ile Leu Leu Arg Trp Met Glu Glu  
 115                      120                      125  
 Met Leu  
 130

<210> 36  
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 <212> DNA  
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<400> 36  
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<210> 37  
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 <212> DNA  
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<400> 37 gcgaggctga tctttct	17
<210> 38 <211> 25 <212> DNA <213> Mus musculus	
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<210> 43

<211> 25

<212> DNA

<213> Mus musculus

<400> 43

attgcagtgt aagggaatac agaga